

Figure 1

Polynucleotide sequence of Human BMP2A gene (gi: 179501 and 4557368) (SEQ ID NO:1)
(Coding sequence: 324 – 1514)

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1  ggggacttct tgaacttgca gggagaataa cttgcgcacc ccactttgcg ccggtgcctt
61  tgccccagcg gagcctgctt cgccatctcc gagccccacc gccctccac tcctcggcct
121 tgccccgacac tgagacgctg ttcccagcgt gaaaagagag actgcgcggc cggcaccg
181 gagaaggagg aggcaaagaa aaggaaacgga cattcgggtcc ttgcgccagg tcctttgacc
241 agagtttttc catgtggacg ctctttcaat ggacgtgtcc ccgcgtgctt cttagacgga
301 ctgcggtctc ctaaaggctg accatgggtgg ccgggacccg ctgtcttcta gcgttgctgc
361 ttccccaggt cctcctgggc ggcgcggctg gcctcgttcc ggagctgggc cgcaggaagt
421 tcgcggcggc gtcgtcgggc cgccctcat cccagccctc tgacgaggtc ctgagcgagt
481 tcgagttgcg gctgctcagc atgttcggcc tgaaacagag acccaccccc agcagggacg
541 ccgtggtgcc ccctacatg ctagacctgt atcgcaggca ctcaggtcag ccgggctcac
601 ccgccccaga ccaccggttg gagagggcag ccagccgagc caacactgtg cgcagcttcc
661 accatgaaga atctttggaa gaactaccag aaacgagtgg gaaaacaacc cggagattct
721 tctttaattt aagttctatc ccacggagg agtttatcac ctcagcagag cttcaggttt
781 tccgagaaca gatgcaagat gctttaggaa acaatagcag tttccatcac cgaattaata
841 tttatgaaat cataaaacct gcaacagcca actcgaaatt ccccgtgacc agacttttgg
901 acaccagggt ggtgaatcag aatgcaagca ggtgggaaag ttttgatgtc acccccgtg
961 tgatgcggtg gactgcacag ggacacgcca accatggatt cgtggtggaa gtggcccact
1021 tggaggagaa acaagggtgc tccaagagac atgttaggat aagcaggctt ttgcaccaag
1081 atgaacacag ctggtcacag ataaggccat tgctagtaac ttttggccat gatggaaaag
1141 ggcatacctt ccacaaaaga gaaaaacgtc aagccaaaca caaacagcgg aaacgcctta
1201 agtccagctg taagagacac cctttgtacg tggacttcag tgacgtgggg tggaatgact
1261 ggattgtggc tccccgggg tatcacgcct tttactgcca cggagaatgc ccttttcctc
1321 tggctgatca tctgaactcc actaatcatg ccattgttca gacgttggtc aactctgtta
1381 actctaagat tcctaaggca tgctgtgtcc cgacagaact cagtgtatc tcgatgctgt
1441 accttgacga gaatgaaaag gttgtattaa agaactatca ggacatggtt gtggagggtt
1501 gtgggtgtcg ctagtacagc aaaattaaat acataaatat atatata
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Figure 2

Polypeptide sequence of human BMP2A (SEQ ID NO:2 = translation of SEQ ID NO:1)

5	1	MVAGTRCLLA	LLLPQVLLGG	AAGLVPELGR	RKFAAASSGR
	41	PSSQPSDEVL	SEFELRLLSM	FGLKQRPTPS	RDAVVPPYML
	81	DLYRRHSGQP	GSPAPDHRLE	RAASRANTVR	SFHHEESLEE
	121	LPETSGKTTR	RFFFNLSIP	TEEFITSAEL	QVFREQMQDA
	161	LGNNSSFHHR	INIYEIIKPA	TANSKFPVTR	LLDTRLVNQN
10	201	ASRWESFDVT	PAVMRWTAGG	HANHGFVVEV	AHLEEKQGVV
	241	KRHVRISRSL	HQDEHSWSQI	RPLLVTFGHD	GKGHPLHKRE
	281	KRQAKHKQRK	RLKSSCKRHP	LYVDFSDVGW	NDWIVAPPGY
	321	HAFYCHGECF	FPLADHLNST	NHAIVQTLVN	SVNSKIPKAC
	361	CVPTELSAIS	MLYLDENEKV	VLKNYQDMVV	EGCGCR

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Figure 3
Results of a loss of function validation experiment

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